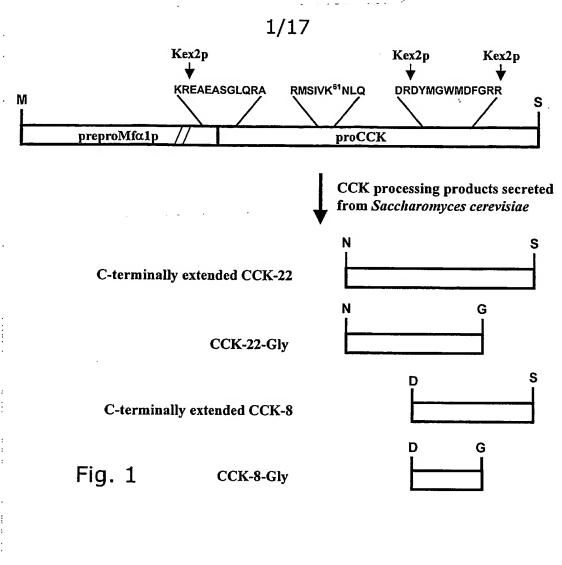
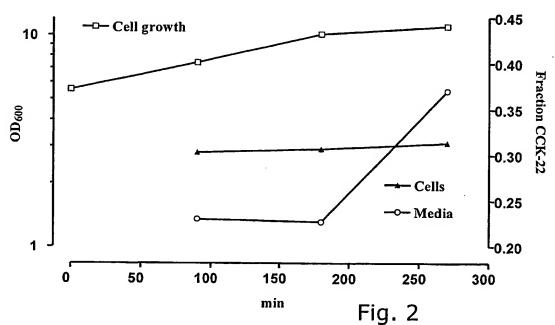
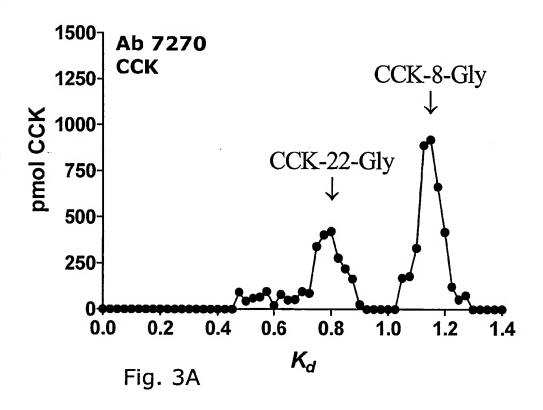
DOCKET NO.: 030307-0256

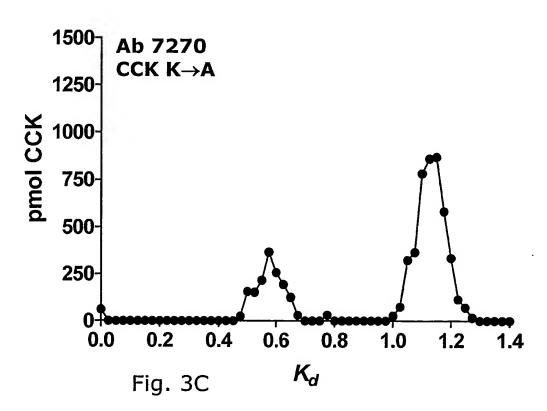
WO 2004/027067





PCT/DK2003/000609



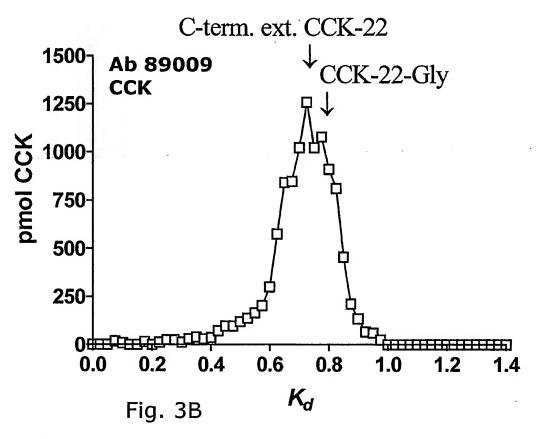


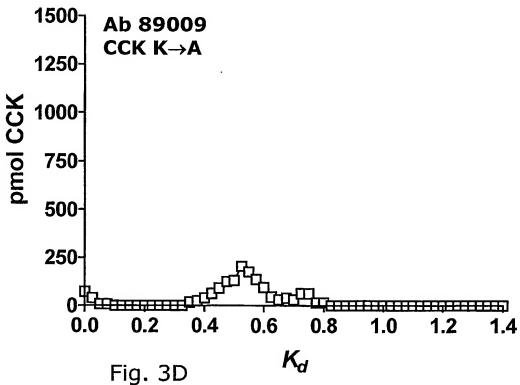
MAP ! OS SAME.

WO 2004/027067

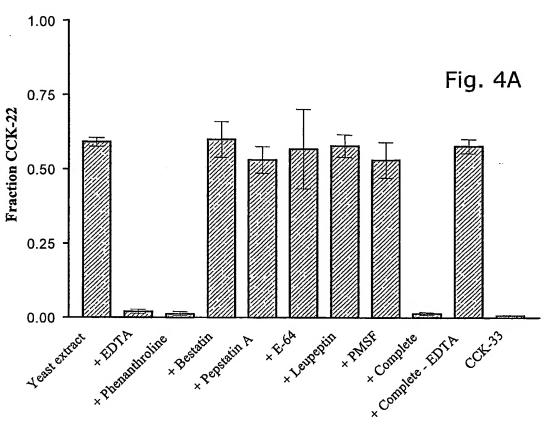
Title: METHODS FOR INCREASING THE PRODUCTION OF A RECOMBINANT POLYPEPTIDE FROM A HOST CELL Inventor(s): Lars JONSON et al. DOCKET NO.: 030307-0256

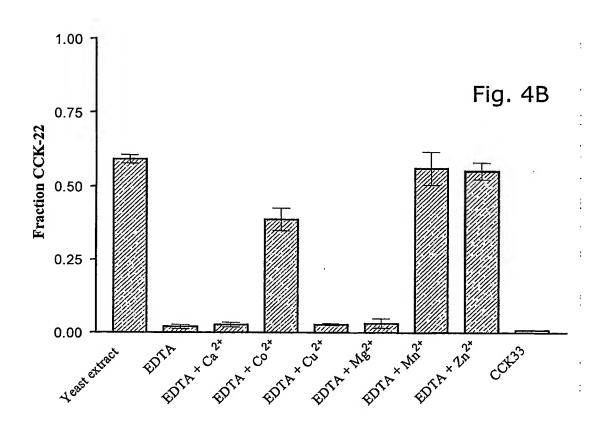
PCT/DK2003/000609







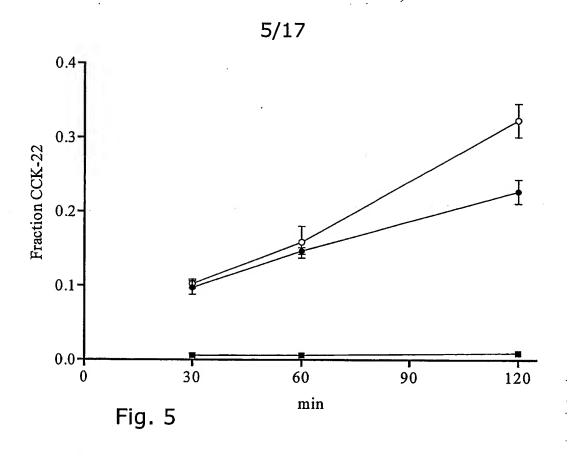


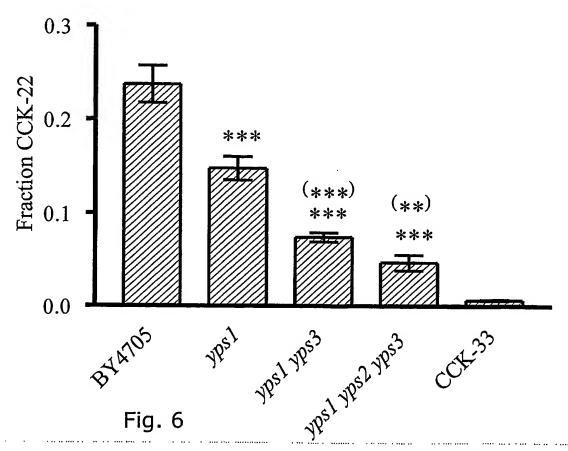


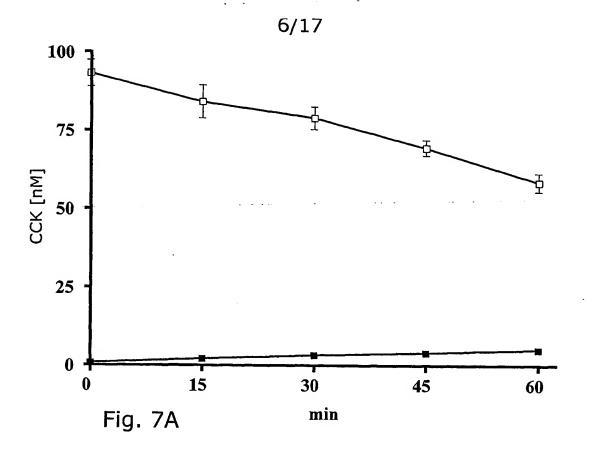
DOCKET NO.: 030307-0256

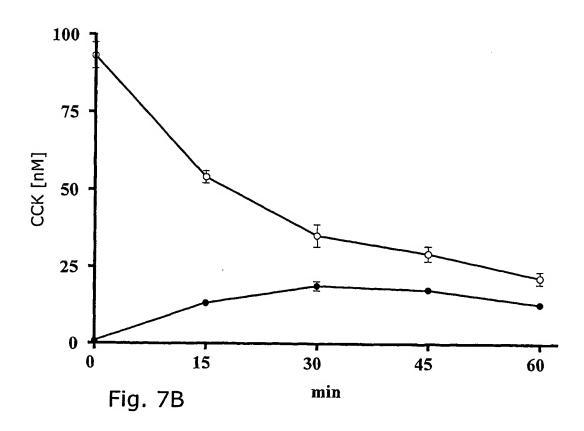
10/528563

WO 2004/027067









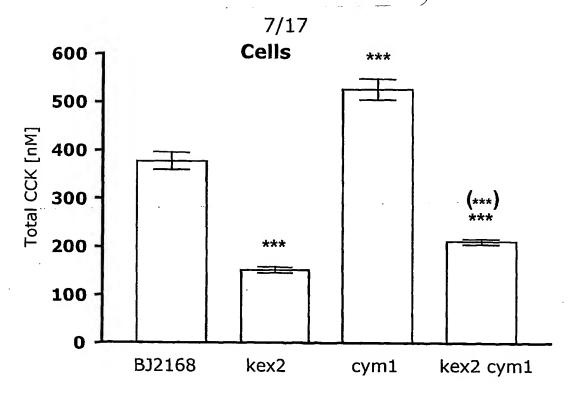


Fig. 8A

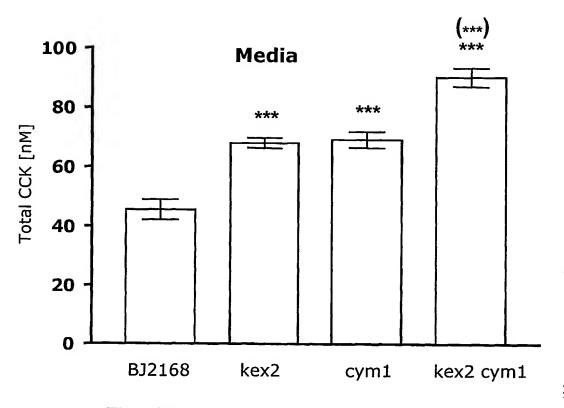


Fig. 8B

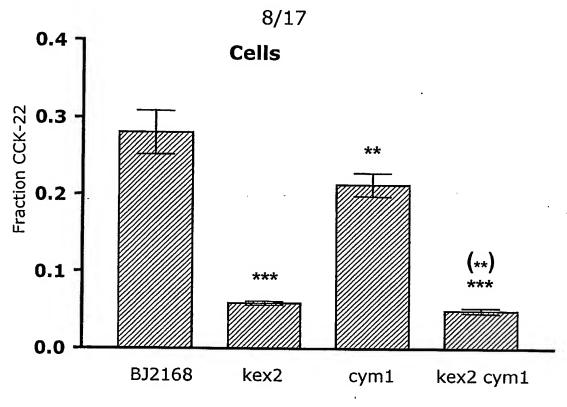


Fig. 8C

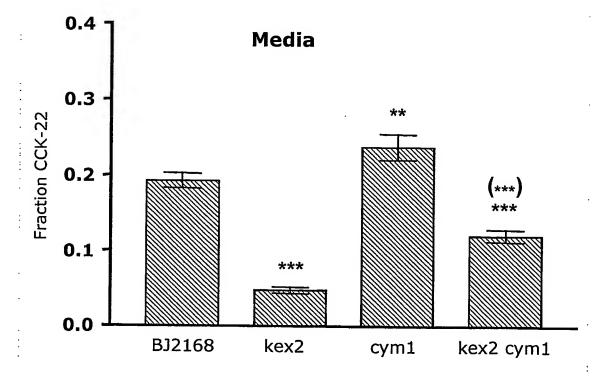


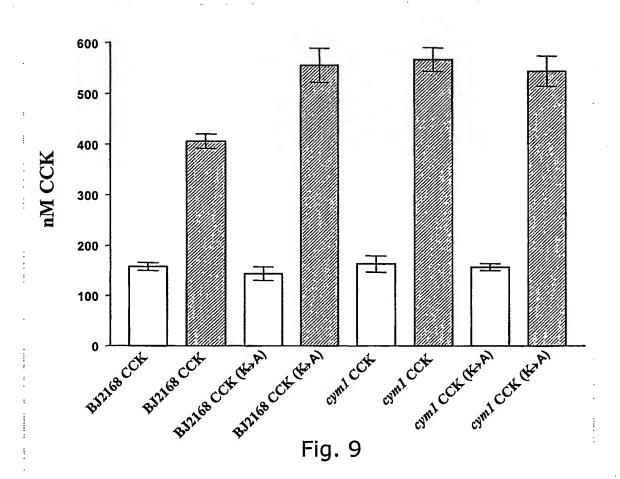
Fig. 8D

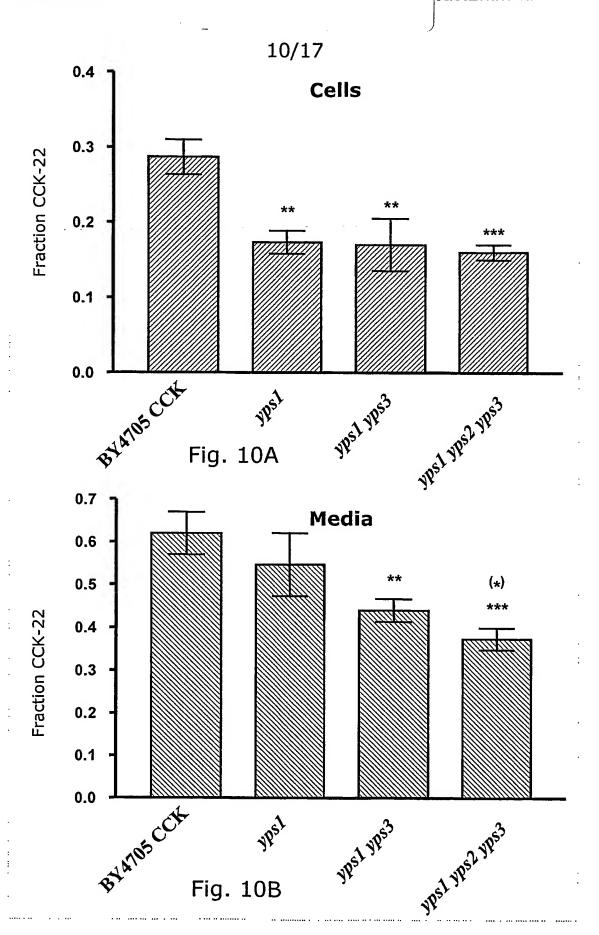
Title: METHODS FOR INCREASING THE PRODUCTION OF A RECOMBINANT POLYPEPTIDE FROM A HOST CELL Inventor(s): Lars IONSON et al. DOCKET NO.: 030307-0256

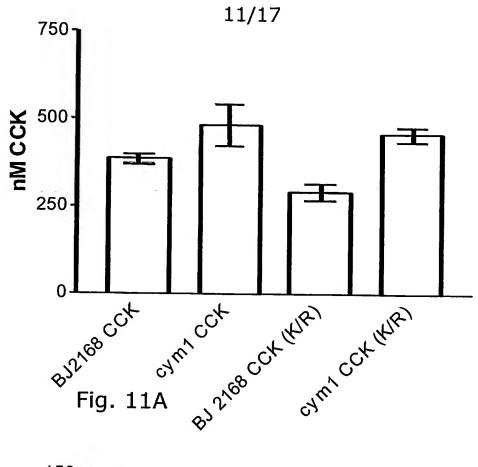
10/528563

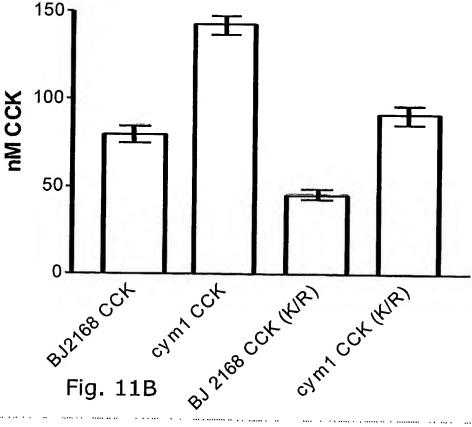
WO 2004/027067

Г/DK2003/000609









PCT/DK2003/000609

	12/17													
SEQ ID	43	44	45	46	47	48	49	20	51	25				
Strain SEQ ID NO:		Ø	മ	щ	8 9 4	æ	æ	æ	A & B	æ				
		2433.5	2036.1	6051.6*	1932.2	1805.0	1508.7	2766.1	4133.9	2488.1				
Molecular mass		2433.3	2036.1	6051.9*	1932.1	1805.2	1509.0	2766.2	4133.8	2488.0				
<u>ProCCK</u>	CCK-61 CCK-58 CCK-39 CCK-22 CCK-8 Prepromíalp //.SGLQRAEEAPRRALRVSQRTDGESRAHLGALLARYIQQARKAPSGRMSIVKNLQNLDPSHRISDRDYMGWMDFGRRSAEEYEYPS	QLRVSQRTDGESRAHLGALLAR	VSQRTDGESRAHLGALLAR	YIQQARKAPSGRMSIVKNLQNLDPSHRISDRDYMGWMDFGRRSAEEYEYPS	YIQQARKAPSGRMSIVK	YIQQARKAPSGRMSIV	NLQNLDPSHRISD	NLQNLDPSHRISDRDYMGWMDFG	NLQNLDPSHRISDRDYMGWMDFGRRSAEEYEYPS	DYMGWNDFGRRSAEBYEYPS				

FIG. 12

DOCKET NO.: 030307-0256

10/528563

WO 2004/027067

CT/DK2003/000609

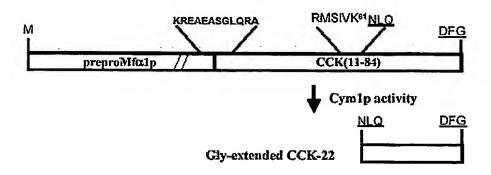


Fig. 13A

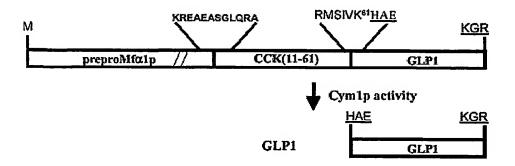


Fig. 13B

CT/DK2003/000609

-	A:	(GA	SATI	TCC	CTT	CAA'	TTT	TTA	CTG	CAG	TTT	TAT	TCG	CAG	CAT	CCT	CCG	CAT:	ľAGO	CTGCT	
;	TA	ACTC	TAA F	AAG	SAA	GTT	AAA	AAT	GAC	GTC.	AAA	ATA	AGC	+ GTC A	GTA	GGA	GGC	GTA/	ATCO	+ SACGA	60
																_		_		A	-
. 61				-+-				+			-+-			+-				+		CGGT	120
	P	V	N	T	- T	T.	E E	D	E	T	A	ĠŦĞ Q	ITT	AAG(P	GCC(A	GAC' E	TTC(A			GCCA G	_
121	TA	CTI	'AGA	TTI	'AG	AAG	GGG	ATT'	TCG	ATG'	TTG	CTG'	TTT?	rgco	CAT	r t r	CCA	ACAC	CAC	TAAA	
. 121	AI	'GAA	TCT	AAA	ATCI	CTC	CCC!	TAA	AGC'	TAC	AAC	GAC	AAA	ACGO	GTA/	AAA	GGT	 CGT(GTG	+ TTTA	180
			D			_								P		-		-	_	N	-
181				-+-				+			-+			+-						GGTA	240
:	N	GCC	CAA L	TAA L	CAA F	ATA I	ATT! N	TAT) T	GAT(T		AAC A			AACO A				TCT E		CCAT V	_
	TC	TTT	GGA	TAA	AAG	SAGA	AGG	CTG	AAG	CTC	ACCO	CGC	rGGG	CAG	CCC	CCGG	STTC	CAGO	CTC	GGAC	
. 241	AG	AAA	CCT.	ATT	TTC	TCI	CCC	SAC'	TCC	GAG:	rgg(SCG/	ACCC	CGTC	CGGG	GCC	CAAC	TCG	GAG	+ CCTG	300
:	s 		D														_		_	D	-
301				-+-			+				-+			-+-			+			GCTG	360
	AA L	E E	TTG(T	CAG S	GCC G	CAA L	TGT Q	CCI E	CGI Q	CGC R	CGTI N	'GG'I H	'AAA' L	CGT Q	CCC G	GTI K	TGA L	CAG S	_	CGAC L	_
261	L E T S G L Q E Q R N H L Q G K L S E L CAGGTGGAGCAGACATCCCTGGAGCCCCTCCAGGAGAGCCCCCGTCCCACAGGTGTCTGG																				
201	GT	CCA	CCT	CGT	CTG	TAG	GGA	CCI	'CGG	GGA	GGI	'CCT	CTC	GGG	GGC	AGG	GTG	TCC	ACA	+ GACC	420
	-	V	-	Q	T 									P				_	V		_
421				-+-			+				+			-+-			+			CACC	480
	K	S	GGC R	E E	CCA V	TCG A	GTG T	GCT E	CCC G					GGC R		TTA M	.CCA V	GGA L	GAT(Y	GTGG T	_
401	СТ	GCGC	GCF	ACC	ACG	AAG	ccc	CAA	GAT	GGT	GCA	AGG	GTC	TGG	CTG	CTT	TGG	GAG	GAA	GATG	
481	GAG	CGCC	CGI	GG:	rgc:	TTC	GGG	GTT	CTA	CCA	CGT	TCC	CAG	ACC	GAC	GAA	ACC	СТС	ርጥጥር	ישבר	540
	Ь	R	Α	P	R	S	P	K	M	V	Q	G	S	G	С	F	G	R	K	M	-
541			ATC	+			+				+			-+-			+		- 59	94	
	D	R R	TAG I	STC S	SAG(GAG(S	GTC: S	ACC G	GGA L	CCC G	GAC C	GTT K	TCA V	CGA(CTC R			AAT:	r –		

DOCKET NO.: 030307-0256 WO 2004/027067

PCT/DK2003/000609

15/17

1	AT	GAG	ATT	TCC	TTC	AAT	TTT	'TAC	TGC	AGT	TTI	TAT.	'CGC	AGC	ATO	CTC	CGC	:ATT	'AGC	TGCT	C 0											
	TA	CTC	TAA	AGG	AAG	TTA	AAA	ATG	ACG	TCA	AAA	TAA	GCG	TCG	TAC	GAC	GCG	TAA	TCG	+ ACGA	60											
	M	R	F	P	S	I	F	T	A	V	L	F	A	A	s	s	A	L	A	A	-											
61	CC.	AGT	CAA	CAC	TAC	AAC	AGA	AGA	TGA	AAC	GGC	ACA	IAAI	TCC	GGC	TGF	AGC	TGI	CAT	CGGT												
-	. GG	TCA	GTT	GTG	ATG	TTG	TCT	TCT	ACT	TTG	CCG	TGI	TTA	AGG	CČC	AC1	TCG	ACA	GTA	GCCA	120											
	P	V	N	T	T	T	E	D	E	T	A	Q	I	P	A	E	A	v	I	G	_											
61 61 121 181 301	TA	TACTTAGATTTAGAAGGGGATTTCGATGTTGCTGTTTTTGCCATTTTCCAACAGCACAAAT																														
	AT	GAA'	rct.	AAA	TCT	TCC	CCT	AAA	GCT	ACA	ACG	ACA	AAA	.CGG	TAA	AAG	GTT	GTC	GTG	+ TTTA	'TA 'TA											
	Y	L	D	L	Е	G	D	F	D	٧	A	V	L	P	F	s	N	s	T	N	-											
1 61 121 181 181 181 181 181 181 181 181	AA	ACGGGTTATTGTTTATAAATACTACTATTGCCAGCATTGCTGCTAAAGAAGAAGAGGGGTA															0.40															
	TT	GCC	CAA!	raa	CAA	ATA'	TTT.	ATG	ATG	ATA	ACG	GTC	GTA	ACG	ACG	ATI	TCT	TCT	TCC	TCCCCAT												
	N	G	L	L	F	I	N	T	T	I	A	s	I	A	A	K	E	E	G	V	_											
241	TC:	TTT(GA:	TAA	AAG	AGA	GGC	TGA.	AGC	TAG	CCC	CAA	GAT	GGT	GCA	AGG	GTC	TGG	CTG	CTTT	200											
	AG	AGAAACCTATTTTCTCTCCGACTTCGATCGGGGTTCTACCACGTTCCCAGACCGACGAAA															300															
	s	L	D	K	R	E	A	E	A	s	P	K	M	v	Q	G	s	G	С	F	-											
801	GGG	GGGAGGAAGATGGACCGGATCAGCTCCTCCAGTGGCCTGGGCTGCAAAGTGCTGAGGCGG															260															
61	CCC	CTC	CTTC	TAC	CCT	GGC	CTA	GTC	GAG	GAG	GTC	ACC	GGA	CCC	GAC	GTT	TCA	CGA	CTC	CGCC	360											
	G	R	K	M	D	R	I	s	s	s	s	G	L	G	С	K	v	L	R	R	_											
61		TA		56																												
		LTA																														
	H	*	-						ı			1 1	D																			

Fig. 14B

CT/DK2003/000609

	A	rgac	SAT'	TTC	CTT	CAA'	TTT'	TTA	CTG	CAG	ттт	TAT'	TCG	CAG	CAT	CCT	CCG	CAT	TAG	CTGCI	1							
1														 ATC	60													
	M	R	F	P	S	I	F	T	A	V	L	F	A	A	s	s	Α	L	A	A	-							
61	CC	CCAGTCAACACTACAACAGAAGATGAAACGGCACAAATTCCGGCTGAAGCTGTCATCGGT															TCGGT											
	GG	TCA	GTT	GT	SATO	STTO	STC	rTC:	rac:	rtt(GCC(GTG:	rtti	AAG	GCC	GAC	TTC	F GAC	AGT	AGCCA	120							
	P	V	N	T	T	T	E	D	E	T	A	Q	I	P	A	E	A	V	I	G	-							
121	TA	TACTTAGATTTAGAAGGGGATTTCGATGTTGCTGTTTTGCCATTTTCCAACAGCACAAAT																										
	AT	GAA	TCI	'AAZ	ATCI	TCC	CCI	'AAZ	AGCI	AC	AAC(SACA	AAA.	ACG(TA	+ 180 AAAGGTTGTCGTGTTTA												
181	Y	L	D	L	E	G	D	F	D	V	A	V	L	P	F	s	N	s	T	N	-							
	AA	AACGGGTTATTGTTTATAAATACTACTATTGCCAGCATTGCTGCTAAAGAAGAAGGGGGTA																										
	TT	TTGCCCAATAACAAATATTTATGATGATAACGGTCGTAACGACGATTTCTTCCCCAT															240											
	N	G	L	L	F	I	N	Т	T	I	A	s	I	A	A	K	Е	E	G	v	_							
241	TC	TCTTTGGATAAAAGAAGCCCCAAGATGGTGCAAGGGTCTGGCTGCTTTGGGAGGAAGATG																										
241	AG	AAA	CCT	-+- ATT	TTC	TTC	GGG	 GTT	CTA	.CCA	.CGI	TCC	CAG	ACC	GAC	GAA	ACC	CTC	CTI	CTAC	300							
	S	L	D	K	R	s	P	K	M	V	Q	G	s	G	С	F	G	R	K	M	_							
301	GA	CCG	GAT	CAG	CTC	CTC	CAG	TGG	CCT	GGG	CTG	CAA	AGT	GCT	GAG	GCG	GCA	TTA	A									
	CT	GGC	CTA	GTC	GAG	GAG	GTC	ACC	GGA	CCC	+ GAC	 GTT	 TCA	~+- CGA	 CTC	CGC	+ CGT	 AAT	– 3 T	54								
	D	R	I	s	s	s	s	G	L	G	С	K	v	${f r}$	R	R	Н	*	_									

Fig. 14C

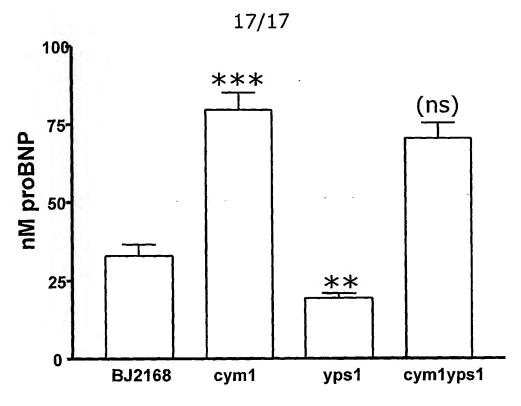


Fig. 15A

